

Resource Summary Report

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ECOGEO 'Omics Training: 3.1 Amplicon Analysis - Mothur

DOI:10.17504/protocols.io.fi4bkgw

Type: Protocol

Proper Citation

Ben Tully 2016. ECOGEO 'Omics Training: 3.1 Amplicon Analysis - Mothur. protocols.io dx.doi.org/10.17504/protocols.io.fi4bkgw

Protocol Information

URL: <https://dx.doi.org/10.17504/protocols.io.fi4bkgw>

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Group: ECOGEO

Summary: mothur is a bioinformatics tool for analyzing 16S rRNA gene sequences, which can be used to ??process data generated by Sanger, PacBio, IonTorrent, 454, and Illumina (MiSeq/HiSeq).

Affiliations: EarthCube Oceanography and Geobiology Environmental 'Omics

Version: 2

Publication Date: 2016

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